

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: <u>10700,971</u>
ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE		
1 _____ Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."	
2 _____ Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.	
3 _____ Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters , instead.	
4 _____ Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.	
5 _____ Variable Length	Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.	
6 _____ PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.	
7 _____ Skipped Sequences (OLD RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO: X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION: SEQ ID NO: X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.	
8 _____ Skipped Sequences (NEW RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000	
9 _____ Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.	
10 _____ Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence	
11 _____ Use of <220>	Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)	
12 _____ PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.	
13 _____ Misuse of n/Xaa	"n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid	



IFWO

RAW SEQUENCE LISTING

DATE: 11/17/2003

PATENT APPLICATION: US/10/700,971

TIME: 09:59:22

Input Set : A:\ISIC0009-101.txt

Output Set: N:\CRF4\11172003\J700971.raw

3 <110> APPLICANT: Manoharan, Muthiah
 4 Baker, Brenda
 5 Eldrup, Ann
 6 Bhat, Balkrishen
 7 Griffey, Richard H.
 8 Swayze, Eric E.
 9 Crooke, Stanley T.
 12 <120> TITLE OF INVENTION: Conjugated Oligomeric Compounds and Their Use in Gene
 13 Modulation
 15 <130> FILE REFERENCE: ISIC-0009-101
 C--> 17 <140> CURRENT APPLICATION NUMBER: US/10/700,971
 C--> 17 <141> CURRENT FILING DATE: 2003-11-04
 17 <150> PRIOR APPLICATION NUMBER: US 10/616,241
 18 <151> PRIOR FILING DATE: 2003-07-09
 20 <150> PRIOR APPLICATION NUMBER: US 60/423,760
 21 <151> PRIOR FILING DATE: 2002-11-05
 23 <150> PRIOR APPLICATION NUMBER: US 10/078,949
 24 <151> PRIOR FILING DATE: 2002-02-20
 26 <150> PRIOR APPLICATION NUMBER: US 09/479,783
 27 <151> PRIOR FILING DATE: 2000-01-07
 29 <150> PRIOR APPLICATION NUMBER: US 08/870,608
 30 <151> PRIOR FILING DATE: 1997-06-06
 32 <150> PRIOR APPLICATION NUMBER: US 08/659,440
 33 <151> PRIOR FILING DATE: 1996-06-06
 36 <160> NUMBER OF SEQ ID NOS: 26
 38 <170> SOFTWARE: PatentIn version 3.2
 40 <210> SEQ ID NO: 1
 41 <211> LENGTH: 16
 42 <212> TYPE: PRT
 43 <213> ORGANISM: Artificial Sequence
 45 <220> FEATURE:
 46 <223> OTHER INFORMATION: (Peptide) insufficient explanation. Give source of
 48 <400> SEQUENCE: 1
 50 Arg Gln Ile Lys Ile Trp Phe Gln Asn Arg Arg Met Lys Trp Lys Lys
 51 1 5 10 15
 55 <210> SEQ ID NO: 2
 56 <211> LENGTH: 13
 57 <212> TYPE: PRT
 58 <213> ORGANISM: Artificial Sequence
 60 <220> FEATURE:
 61 <223> OTHER INFORMATION: (Peptide)
 63 <400> SEQUENCE: 2
 65 Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln

pp 1-5

Give source of
 genetic
 material.
 (see item 11
 on Error
 Summary sheet)

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71 <212> TYPE: PRT
72 <213> ORGANISM: Artificial Sequence
74 <220> FEATURE:
75 <223> OTHER INFORMATION: Peptide
77 <400> SEQUENCE: 3
79 Gly Trp Thr Leu Asn Ser Ala Gly Tyr Leu Leu Gly Pro Ile Asn Leu
80 1          5          10          15
82 Lys Ala Leu Ala Ala Leu Ala Lys Lys Ile Leu
83          20          25
86 <210> SEQ ID NO: 4
87 <211> LENGTH: 34
88 <212> TYPE: PRT
89 <213> ORGANISM: Artificial Sequence
91 <220> FEATURE:
92 <223> OTHER INFORMATION: Peptide - HSV VP22
94 <400> SEQUENCE: 4
96 Asp Ala Ala Thr Ala Thr Arg Gly Arg Ser Ala Ala Ser Arg Pro Thr
97 1          5          10          15
99 Glu Arg Pro Arg Ala Pro Ala Arg Ser Ala Ser Arg Pro Arg Arg Pro
100          20          25          30
102 Val Glu
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106 <211> LENGTH: 18
107 <212> TYPE: PRT
108 <213> ORGANISM: Artificial Sequence
110 <220> FEATURE:
111 <223> OTHER INFORMATION: Peptide
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116 1          5          10          15
118 Leu Ala
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122 <211> LENGTH: 27
123 <212> TYPE: PRT
124 <213> ORGANISM: Artificial Sequence
126 <220> FEATURE:
127 <223> OTHER INFORMATION: Peptide
129 <400> SEQUENCE: 6
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132 1          5          10          15
134 Ala Trp Ser Gln Pro Lys Lys Lys Arg Lys Val
135          20          25
138 <210> SEQ ID NO: 7
139 <211> LENGTH: 16
140 <212> TYPE: PRT
141 <213> ORGANISM: Artificial Sequence

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OK

RAW SEQUENCE LISTING

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143 <220> FEATURE:
144 <223> OTHER INFORMATION: Peptide
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148 Ala Ala Val Ala Leu Leu Pro Ala Val Leu Leu Ala Leu Leu Ala Pro
149 1 5 10 15
152 <210> SEQ ID NO: 8
153 <211> LENGTH: 7
154 <212> TYPE: PRT
155 <213> ORGANISM: Artificial Sequence
157 <220> FEATURE:
158 <223> OTHER INFORMATION: Peptide
160 <400> SEQUENCE: 8
162 Pro Lys Lys Lys Arg Lys Val
163 1 5
166 <210> SEQ ID NO: 9
167 <211> LENGTH: 4
168 <212> TYPE: PRT
169 <213> ORGANISM: Artificial Sequence
171 <220> FEATURE:
172 <223> OTHER INFORMATION: Peptide
174 <400> SEQUENCE: 9
176 Met Leu Phe Tyr
177 1
180 <210> SEQ ID NO: 10
181 <211> LENGTH: 15
182 <212> TYPE: PRT
183 <213> ORGANISM: Artificial Sequence
185 <220> FEATURE:
186 <223> OTHER INFORMATION: Peptide - FXR2P
188 <400> SEQUENCE: 10
190 Pro Gln Arg Arg Asn Arg Ser Arg Arg Arg Arg Phe Arg Gly Gln
191 1 5 10 15
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196 <212> TYPE: PRT
197 <213> ORGANISM: Artificial Sequence
199 <220> FEATURE:
200 <223> OTHER INFORMATION: Peptide
202 <400> SEQUENCE: 11
204 Ile Met Arg Arg Arg Gly Leu
205 1 5
208 <210> SEQ ID NO: 12
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210 <212> TYPE: PRT
211 <213> ORGANISM: Artificial Sequence
213 <220> FEATURE:
214 <223> OTHER INFORMATION: Peptide
216 <400> SEQUENCE: 12
218 Leu Gln Leu Pro Pro Leu Glu Arg Leu Thr Leu

OK

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219 1 5 10
222 <210> SEQ ID NO: 13
223 <211> LENGTH: 11
224 <212> TYPE: PRT
225 <213> ORGANISM: Artificial Sequence
227 <220> FEATURE:
228 <223> OTHER INFORMATION: Peptide
230 <400> SEQUENCE: 13
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233 1 5 10
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237 <211> LENGTH: 11
238 <212> TYPE: PRT
239 <213> ORGANISM: Artificial Sequence
241 <220> FEATURE:
242 <223> OTHER INFORMATION: Peptide
244 <400> SEQUENCE: 14
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247 1 5 10
250 <210> SEQ ID NO: 15
251 <211> LENGTH: 12
252 <212> TYPE: PRT
253 <213> ORGANISM: Artificial Sequence
255 <220> FEATURE:
256 <223> OTHER INFORMATION: Peptide
258 <400> SEQUENCE: 15
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261 1 5 10
264 <210> SEQ ID NO: 16
265 <211> LENGTH: 7
266 <212> TYPE: PRT
267 <213> ORGANISM: Artificial Sequence
269 <220> FEATURE:
270 <223> OTHER INFORMATION: Peptide
272 <400> SEQUENCE: 16
274 Pro Lys Lys Lys Arg Lys Val
275 1 5
278 <210> SEQ ID NO: 17
279 <211> LENGTH: 13
280 <212> TYPE: PRT
281 <213> ORGANISM: Artificial Sequence
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284 <223> OTHER INFORMATION: Peptide
286 <400> SEQUENCE: 17
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289 1 5 10
292 <210> SEQ ID NO: 18
293 <211> LENGTH: 4
294 <212> TYPE: PRT

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Input Set : A:\ISIC0009-101.txt

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 297 <220> FEATURE:
 298 <223> OTHER INFORMATION: Peptide
 300 <400> SEQUENCE: 18
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 303 1
 305 <210> SEQ ID NO: 19
 306 <211> LENGTH: 21
 307 <212> TYPE: DNA
 308 <213> ORGANISM: Artificial Sequence
 310 <220> FEATURE:
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 314 cgagaggcgg acgggaccgt t 21
 317 <210> SEQ ID NO: 20
 318 <211> LENGTH: 21
 319 <212> TYPE: DNA
 320 <213> ORGANISM: Artificial Sequence
 322 <220> FEATURE:
 323 <223> OTHER INFORMATION: oligonucleotide
 325 <400> SEQUENCE: 20
 326 ttgctctccg cctgccctgg c 21
 329 <210> SEQ ID NO: 21
 330 <211> LENGTH: 21
 331 <212> TYPE: DNA
 332 <213> ORGANISM: Artificial Sequence
 334 <220> FEATURE:
 335 <223> OTHER INFORMATION: oligonucleotide - cRaf targeter
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 338 augcauguca caggcgggat t 21
 341 <210> SEQ ID NO: 22
 342 <211> LENGTH: 21
 343 <212> TYPE: DNA
 344 <213> ORGANISM: Artificial Sequence
 346 <220> FEATURE:
 347 <223> OTHER INFORMATION: oligonucleotide - cRaf targeter
 349 <400> SEQUENCE: 22
 350 ucccgccugu gacaugcaut t 21
 353 <210> SEQ ID NO: 23
 354 <211> LENGTH: 18
 355 <212> TYPE: DNA
 356 <213> ORGANISM: Artificial Sequence
 358 <220> FEATURE:
 359 <223> OTHER INFORMATION: antisense oligonucleotide
 361 <400> SEQUENCE: 23
 362 tgggagccat agcgaggc 18
 365 <210> SEQ ID NO: 24
 366 <211> LENGTH: 20
 367 <212> TYPE: DNA

*Please correct this type of error
in subsequent sequences*

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/700,971

DATE: 11/17/2003

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Input Set : A:\ISIC0009-101.txt

Output Set: N:\CRF4\11172003\J700971.raw

L:17 M:270 C: Current Application Number differs, Replaced Current Application No

L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date